


IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

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Serial No.: Not yet assigned)
) I hereby certify that this paper and
Filed: Herewith) the documents referred to as
) enclosed herewith are being
For: Flt4 (VEGFR-3) as a Target) deposited with the United States
for Tumor Imaging and Anti-Tumor) Postal Service as "EXPRESS MAIL
Therapy) POST OFFICE TO ADDRESSEE"
) Service under 37 C.F.R. §1.10 on
Group Art Unit: Not yet assigned) the date indicated below and is
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) Washington, D.C. 20231 on this
) date:
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) January 19, 2001
)
) 
) David A. Gass

STATEMENT PURSUANT TO 37 C.F.R. § 1.921(f)

Commissioner for Patents
Box Patent Application
Washington, D.C. 20231

Sir:


I hereby state that the content of the paper and computer readable forms of the sequence listing that is part of the above-identified application and that are filed herewith are the same.

Respectfully submitted,

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Reg. No: 38,153

SEQUENCE LISTING

<110> Alitalo, Kari
Kaipainen, Arja
Valtola, Reija
Jussila, Lotta

<120> Flt4 (VEGFR-3) as a Target for Tumor Imaging and Anti-Tumor Therapy

<130> 28113/34891

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<150> 08/901,710

<151> 1997-07-28

<150> 08/340,011

<151> 1994-11-14

<150> 08/257,754

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- 12 -

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Gly	Asp	Ser	Leu	Ser	Ile	Ser	Cys	Arg	Gly	Gln	His	Pro	Leu	Glu	Trp	
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Ala	Trp	Pro	Gly	Ala	Gln	Glu	Ala	Pro	Ala	Thr	Gly	Asp	Lys	Asp	Ser	
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Tyr	Cys	Lys	Val	Leu	Leu	Leu	His	Glu	Val	His	Ala	Asn	Asp	Thr	Gly	
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Ser	Tyr	Val	Cys	Tyr	Tyr	Lys	Tyr	Ile	Lys	Ala	Arg	Ile	Glu	Gly	Thr	
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Thr	Ala	Ala	Ser	Ser	Tyr	Val	Phe	Val	Arg	Asp	Phe	Glu	Gln	Pro	Phe	
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Arg	Arg	Gly	Met	Leu	Val	Ser	Thr	Pro	Leu	Leu	His	Asp	Ala	Leu	Tyr	
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Val Ser Gln His Asp Leu Gly Ser Tyr Val Cys Lys Ala Asn Asn Gly	
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Glu Gly Ser Phe Ser Gln Val Ser Thr Met Ala Leu His Ile Ala Gln	
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Ala Asp Ala Glu Asp Ser Pro Pro Ser Leu Gln Arg His Ser Leu Ala
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Ala Glu Thr Arg Gly Ser Ser Arg Met Lys Thr Phe Glu Glu Phe Pro
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Gly Met Val Leu Ala Ser Glu Glu Phe Glu Gln Ile Glu Ser Arg His
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Thr Phe Phe Thr Asp Asn Ser Tyr
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Asn	Ile	Thr	Glu	Glu	Ser	His	Val	Ile	Asp	Thr	Gly	Asp	Ser	Leu	Ser	
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Ile	Ser	Cys	Arg	Gly	Gln	His	Pro	Leu	Glu	Trp	Ala	Trp	Pro	Gly	Ala	
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Leu Asn Leu Ser Thr Leu His Asp Ala His Gly Asn Pro Leu Leu Leu
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Asp Cys Lys Asn Val His Leu Phe Ala Thr Pro Leu Ala Ala Ser Leu
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Pro Arg Val Ala Pro Glu His Glu Gly His Tyr Val Cys Glu Val Gln
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Asp Arg Arg Ser His Asp Lys His Cys His Lys Lys Tyr Leu Ser Val
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Val Asn Val Ser Asp Ser Leu Glu Met Gln Cys Leu Val Ala Gly Ala
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His Ala Pro Ser Ile Val Trp Tyr Lys Asp Glu Arg Leu Leu Glu Glu
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Lys Ser Gly Val Asp Leu Ala Asp Ser Asn Gln Lys Leu Ser Ile Gln
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Arg Val Arg Glu Glu Asp Ala Gly Arg Tyr Leu Cys Ser Val Cys Asn
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Ala Lys Gly Cys Val Asn Ser Ser Ala Ser Val Ala Val Glu Gly Ser
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Glu Asp Lys Gly Ser Met Glu Ile Val Ile Leu Val Gly Thr Gly Val
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Ile Ala Val Phe Phe Trp Val Leu Leu Leu Leu Ile Phe Cys Asn Met
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Arg Arg Pro Ala His Ala Asp Ile Lys Thr Gly Tyr Leu Ser Ile Ile
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Cys Lys Tyr Leu Ala Val Pro Thr Ser Lys Lys Lys Glu Thr Glu Ser
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Ala	Ile	Tyr	Ile	Phe	Ile	Ser	Asp	Thr	Gly	Arg	Pro	Phe	Val	Glu	Met
130						135					140				
Tyr	Ser	Glu	Ile	Pro	Glu	Ile	Ile	His	Met	Thr	Glu	Gly	Arg	Glu	Leu
145					150					155					160
Val	Ile	Pro	Cys	Arg	Val	Thr	Ser	Pro	Asn	Ile	Thr	Val	Thr	Leu	Lys
				165					170					175	
Lys	Phe	Pro	Leu	Asp	Thr	Leu	Ile	Pro	Asp	Gly	Lys	Arg	Ile	Ile	Trp
			180					185					190		
Asp	Ser	Arg	Lys	Gly	Phe	Ile	Ile	Ser	Asn	Ala	Thr	Tyr	Lys	Glu	Ile
		195					200					205			
Gly	Leu	Leu	Thr	Cys	Glu	Ala	Thr	Val	Asn	Gly	His	Leu	Tyr	Lys	Thr
	210					215					220				
Asn	Asn	Tyr	Leu	Thr	His	Arg	Gln	Thr	Asn	Thr	Ile	Ile	Asp	Val	Gln
225					230					235					240
Ile	Ser	Thr	Pro	Arg	Pro	Val	Lys	Leu	Leu	Arg	Gly	His	Thr	Leu	Val
				245					250					255	
Leu	Asn	Cys	Thr	Ala	Thr	Thr	Pro	Leu	Asn	Thr	Arg	Val	Gln	Met	Thr
			260					265					270		
Trp	Ser	Tyr	Pro	Asp	Asn	Asn	Asn	Glu	Lys	Asn	Lys	Arg	Ala	Ser	Val
		275				280						285			
Arg	Arg	Arg	Ile	Asp	Gln	Ser	Asn	Ser	His	Ala	Asn	Ile	Phe	Tyr	Ser
	290					295					300				
Val	Leu	Thr	Ile	Asp	Lys	Met	Gln	Asn	Lys	Asp	Lys	Gly	Leu	Tyr	Thr
305					310					315					320
Cys	Arg	Val	Arg	Ser	Gly	Pro	Ser	Phe	Lys	Ser	Val	Asn	Thr	Ser	Val
				325					330					335	
His	Ile	Tyr	Asp	Lys	Ala	Phe	Ile	Thr	Val	Lys	His	Arg	Lys	Gln	Gln
			340					345					350		
Val	Leu	Glu	Thr	Val	Ala	Gly	Lys	Arg	Ser	Tyr	Arg	Leu	Ser	Met	Lys
		355					360					365			
Val	Lys	Ala	Phe	Pro	Ser	Pro	Glu	Val	Val	Trp	Leu	Lys	Asp	Gly	Leu
	370					375					380				
Pro	Ala	Thr	Glu	Lys	Ser	Ala	Arg	Tyr	Leu	Thr	Arg	Gly	Tyr	Ser	Leu
385					390					395					400
Ile	Ile	Lys	Asp	Val	Thr	Glu	Glu	Asp	Ala	Gly	Asn	Tyr	Thr	Ile	Leu
				405					410					415	
Leu	Ser	Ile	Lys	Gln	Ser	Asn	Val	Phe	Lys	Asn	Leu	Thr	Ala	Thr	Leu
			420					425					430		
Ile	Val	Asn	Val	Lys	Pro	Gln	Ile	Tyr	Glu	Lys	Ala	Val	Ser	Ser	Phe
		435					440					445			

Pro 450	Asp	Pro	Ala	Leu	Tyr	Pro 455	Leu	Gly	Ser	Arg	Gln 460	Ile	Leu	Thr	Cys
Thr 465	Ala	Tyr	Gly	Ile	Pro 470	Gln	Pro	Asn	Thr	Ile 475	Lys	Trp	Phe	Trp	His 480
Pro	Cys	Asn	His	Asn 485	His	Ser	Glu	Ala	Arg 490	Cys	Asp	Phe	Cys	Ser 495	Asn
Asn	Glu	Glu	Ser 500	Phe	Ile	Leu	Asp 505	Asn	Asn	Asn	Asn	Asn 510	Asn	Asn	Ala
Asp	Ser	Asn 515	Met	Gly	Asn	Arg	Ile 520	Glu	Ser	Ile	Thr	Gln 525	Arg	Met	Ala
Ile 530	Ile	Glu	Gly	Lys	Asn 535	Lys	Met	Ala	Ser	Thr	Leu 540	Val	Val	Ala	Asp
Ser 545	Arg	Ile	Ser	Gly	Ile 550	Tyr	Ile	Cys	Ile	Ala 555	Ser	Asn	Lys	Val	Gly 560
Thr	Val	Gly	Arg	Asn 565	Ile	Ser	Phe	Tyr	Ile 570	Thr	Asp	Val	Pro	Asn 575	Gly
Phe	His	Val	Asn 580	Leu	Glu	Lys	Met	Pro 585	Thr	Asn	Asn	Glu	Gly 590	Glu	Asp
Leu	Lys	Leu 595	Ser	Cys	Thr	Val	Asn 600	Lys	Phe	Leu	Tyr	Arg 605	Asp	Val	Thr
Trp	Ile 610	Leu	Leu	Asn	Asn 615	Asn	Asn	Asn	Asn	Asn	Asn 620	Asn	Asn	Asn	Asn
Asn 625	Asn	Asn	Asn	Asn	Arg 630	Thr	Val	Asn	Asn	Arg 635	Thr	Met	His	Tyr	Ser 640
Ile	Ser	Lys	Gln	Lys 645	Met	Ala	Ile	Thr	Lys 650	Glu	His	Ser	Ile	Thr 655	Leu
Asn	Leu	Thr	Ile 660	Met	Asn	Val	Ser	Leu 665	Gln	Asp	Ser	Gly	Thr 670	Tyr	Ala
Cys	Arg	Ala 675	Arg	Asn	Val	Tyr	Thr 680	Gly	Glu	Glu	Ile	Leu 685	Gln	Lys	Lys
Glu	Ile 690	Thr	Ile	Arg	Asp	Gln 695	Glu	Ala	Pro	Tyr	Leu 700	Leu	Arg	Asn	Leu
Ser 705	Asp	His	Thr	Val	Ala 710	Ile	Ser	Ser	Ser	Thr 715	Thr	Leu	Asp	Cys	His 720
Ala	Asn	Gly	Val	Pro 725	Glu	Pro	Gln	Ile	Thr 730	Trp	Phe	Lys	Asn	Asn 735	His
Lys	Ile	Gln	Gln 740	Glu	Pro	Gly	Ile	Ile 745	Leu	Gly	Pro	Gly	Ser 750	Ser	Thr
Leu	Phe	Ile 755	Glu	Arg	Val	Thr	Glu 760	Glu	Asp	Glu	Gly	Val 765	Tyr	His	Cys

Lys	Ala	Thr	Asn	Gln	Lys	Gly	Ser	Val	Glu	Ser	Ser	Ala	Tyr	Leu	Thr
770						775					780				
Val	Gln	Gly	Thr	Ser	Asp	Lys	Ser	Asn	Leu	Glu	Leu	Ile	Thr	Leu	Thr
785					790					795					800
Cys	Thr	Cys	Val	Ala	Ala	Thr	Leu	Phe	Trp	Leu	Leu	Leu	Thr	Leu	Leu
				805					810					815	
Ile	Arg	Lys	Met	Lys	Arg	Ser	Ser	Asn	Ser	Glu	Ile	Lys	Thr	Asp	Tyr
			820					825					830		
Leu	Ser	Ile	Ile	Met	Asp	Pro	Asp	Glu	Val	Pro	Leu	Asp	Glu	Gln	Cys
		835					840					845			
Glu	Arg	Leu	Pro	Tyr	Asp	Ala	Ser	Lys	Trp	Glu	Phe	Ala	Arg	Glu	Arg
850						855					860				
Leu	Lys	Leu	Gly	Lys	Ser	Leu	Gly	Arg	Gly	Ala	Phe	Gly	Lys	Val	Val
865					870					875					880
Gln	Ala	Ser	Ala	Phe	Gly	Ile	Lys	Lys	Ser	Pro	Thr	Cys	Arg	Thr	Val
				885					890					895	
Ala	Val	Lys	Met	Leu	Lys	Glu	Gly	Ala	Thr	Ala	Ser	Glu	Tyr	Lys	Ala
			900					905					910		
Leu	Met	Thr	Glu	Leu	Lys	Ile	Leu	Thr	His	Ile	Gly	His	His	Leu	Asn
		915					920					925			
Val	Val	Asn	Leu	Leu	Gly	Ala	Cys	Thr	Lys	Gln	Gly	Gly	Pro	Leu	Met
930						935					940				
Val	Ile	Val	Glu	Tyr	Cys	Lys	Tyr	Gly	Asn	Leu	Ser	Asn	Tyr	Leu	Lys
945					950					955					960
Ser	Lys	Arg	Asp	Leu	Phe	Phe	Leu	Asn	Lys	Asp	Ala	Ala	Leu	His	Met
				965					970					975	
Glu	Pro	Lys	Lys	Glu	Lys	Met	Glu	Pro	Gly	Leu	Glu	Gln	Gly	Lys	Lys
			980					985					990		
Pro	Arg	Leu	Asp	Ser	Val	Thr	Ser	Ser	Glu	Ser	Phe	Ala	Ser	Ser	Gly
		995					1000					1005			
Phe	Gln	Glu	Asp	Lys	Ser	Leu	Ser	Asp	Val	Glu	Glu	Glu	Glu	Asp	Ser
1010						1015					1020				
Asp	Gly	Phe	Tyr	Lys	Glu	Pro	Ile	Thr	Met	Glu	Asp	Leu	Ile	Ser	Tyr
1025				1030						1035					1040
Ser	Phe	Gln	Val	Ala	Arg	Gly	Met	Glu	Phe	Leu	Ser	Ser	Arg	Lys	Cys
			1045						1050					1055	
Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Leu	Ser	Glu	Asn	Asn
			1060					1065					1070		
Val	Val	Lys	Ile	Cys	Asp	Phe	Gly	Leu	Ala	Arg	Asp	Ile	Tyr	Lys	Asn
	1075						1080					1085			

Pro Asp Tyr Val Arg Lys Gly Asp Thr Arg Leu Pro Leu Lys Trp Met
 1090 1095 1100

Ala Pro Glu Ser Ile Phe Asp Lys Ile Tyr Ser Thr Lys Ser Asp Val
 1105 1110 1115 1120

Trp Ser Tyr Gly Val Leu Leu Trp Glu Ile Phe Ser Leu Gly Gly Ser
 1125 1130 1135

Pro Tyr Pro Gly Val Gln Met Asp Glu Asp Phe Cys Ser Arg Leu Arg
 1140 1145 1150

Glu Gly Met Arg Met Arg Ala Pro Glu Tyr Ser Thr Pro Glu Ile Tyr
 1155 1160 1165

Gln Ile Met Leu Asp Cys Trp His Arg Asp Pro Lys Glu Arg Pro Arg
 1170 1175 1180

Phe Ala Glu Leu Val Glu Lys Leu Gly Asp Leu Leu Gln Ala Asn Val
 1185 1190 1195 1200

Gln Gln Asp Gly Lys Asp Tyr Ile Pro Ile Asn Ala Ile Leu Thr Gly
 1205 1210 1215

Asn Ser Gly Phe Thr Tyr Ser Thr Pro Ala Phe Ser Glu Asp Phe Phe
 1220 1225 1230

Lys Glu Ser Ile Ser Ala Pro Lys Phe Asn Ser Gly Ser Ser Asp Asp
 1235 1240 1245

Val Arg Tyr Val Asn Ala Phe Lys Phe Met Ser Leu Glu Arg Ile Lys
 1250 1255 1260

Thr Phe Glu Glu Leu Leu Pro Asn Ala Thr Ser Met Phe Asp Asp Tyr
 1265 1270 1275 1280

Gln Gly Asp Ser Ser Thr Leu Leu Ala Ser Pro Met Leu Lys Arg Phe
 1285 1290 1295

Thr Trp Thr Asp Ser Lys Pro Lys Ala Ser Leu Lys Ile Glu Val
 1300 1305 1310

<210> 6
 <211> 6
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> The amino acid at positions 1 and 2 each are
 selected independently from the group consisting
 of aspartic acid and glutamic acid.

<220>
 <223> The amino acid at position 4 is independently
 selected from the group consisting of methionine
 and valine.

096544 0190

<220>
<223> Description of Artificial Sequence:
oligonucleotide probe

<400> 10
ccatcgatgg atcctacctg aagccgcttt ctt 33

<210> 11
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide probe

<400> 11
cccaagcttg gatccaagtg gctactccat gacc 34

<210> 12
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide probe

<400> 12
gttgccctgtg atgtgcacca 20

<210> 13
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide probe

<400> 13
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<210> 14
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide probe

<400> 14
cgcggatccc tagtgatggt gatggtgatg totaccttcg atcatgctgc ccttatactc 60

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<220>
<223> Description of Artificial Sequence:
      oligonucleotide probe
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21

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<220>
<223> Description of Artificial Sequence:
      oligonucleotide probe
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44

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Met	His	Leu	Leu	Gly	Phe	Phe	Ser	Val	Ala	Cys	Ser	Leu	Leu	Ala	Ala
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Ala	Leu	Leu	Pro	Gly	Pro	Arg	Glu	Ala	Pro	Ala	Ala	Ala	Ala	Ala	Phe
			20					25					30		
Glu	Ser	Gly	Leu	Asp	Leu	Ser	Asp	Ala	Glu	Pro	Asp	Ala	Gly	Glu	Ala
		35					40					45			
Thr	Ala	Tyr	Ala	Ser	Lys	Asp	Leu	Glu	Glu	Gln	Leu	Arg	Ser	Val	Ser
	50					55					60				
Ser	Val	Asp	Glu	Leu	Met	Thr	Val	Leu	Tyr	Pro	Glu	Tyr	Trp	Lys	Met
65					70					75					80
Tyr	Lys	Cys	Gln	Leu	Arg	Lys	Gly	Gly	Trp	Gln	His	Asn	Arg	Glu	Gln
				85					90					95	
Ala	Asn	Leu	Asn	Ser	Arg	Thr	Glu	Glu	Thr	Ile	Lys	Phe	Ala	Ala	Ala
			100					105					110		
His	Tyr	Asn	Thr	Glu	Ile	Leu	Lys	Ser	Ile	Asp	Asn	Glu	Trp	Arg	Lys
		115					120					125			
Thr	Gln	Cys	Met	Pro	Arg	Glu	Val	Cys	Ile	Asp	Val	Gly	Lys	Glu	Phe
	130					135					140				
Gly	Val	Ala	Thr	Asn	Thr	Phe	Phe	Lys	Pro	Pro	Cys	Val	Ser	Val	Tyr
145					150					155					160
Arg	Cys	Gly	Gly	Cys	Cys	Asn	Ser	Glu	Gly	Leu	Gln	Cys	Met	Asn	Thr
				165					170					175	
Ser	Thr	Ser	Tyr	Leu	Ser	Lys	Thr	Leu	Phe	Glu	Ile	Thr	Val	Pro	Leu
			180					185					190		

Ser Gln Gly Pro Lys Pro Val Thr Ile Ser Phe Ala Asn His Thr Ser
195 200 205

Cys Arg Cys Met Ser Lys Leu Asp Val Tyr Arg Gln Val His Ser Ile
210 215 220

Ile Arg Arg Ser Leu Pro Ala Thr Leu Pro Gln Cys Gln Ala Ala Asn
225 230 235 240

Lys Thr Cys Pro Thr Asn Tyr Met Trp Asn Asn His Ile Cys Arg Cys
245 250 255

Leu Ala Gln Glu Asp Phe Met Phe Ser Ser Asp Ala Gly Asp Asp Ser
260 265 270

Thr Asp Gly Phe His Asp Ile Cys Gly Pro Asn Lys Glu Leu Asp Glu
275 280 285

Glu Thr Cys Gln Cys Val Cys Arg Ala Gly Leu Arg Pro Ala Ser Cys
290 295 300

Gly Pro His Lys Glu Leu Asp Arg Asn Ser Cys Gln Cys Val Cys Lys
305 310 315 320

Asn Lys Leu Phe Pro Ser Gln Cys Gly Ala Asn Arg Glu Phe Asp Glu
325 330 335

Asn Thr Cys Gln Cys Val Cys Lys Arg Thr Cys Pro Arg Asn Gln Pro
340 345 350

Leu Asn Pro Gly Lys Cys Ala Cys Glu Cys Thr Glu Ser Pro Gln Lys
355 360 365

Cys Leu Leu Lys Gly Lys Lys Phe His His Gln Thr Cys Ser Cys Tyr
370 375 380

Arg Arg Pro Cys Thr Asn Arg Gln Lys Ala Cys Glu Pro Gly Phe Ser
385 390 395 400

Tyr Ser Glu Glu Val Cys Arg Cys Val Pro Ser Tyr Trp Lys Arg Pro
405 410 415

Gln Met Ser

<210> 22
<211> 354
<212> PRT
<213> Homo sapiens

<400> 22
Met Tyr Arg Glu Trp Val Val Val Asn Val Phe Met Met Leu Tyr Val
1 5 10 15

Gln Leu Val Gln Gly Ser Ser Asn Glu His Gly Pro Val Lys Arg Ser
20 25 30

Ser	Gln	Ser	Thr	Leu	Glu	Arg	Ser	Glu	Gln	Gln	Ile	Arg	Ala	Ala	Ser
		35					40					45			
Ser	Leu	Glu	Glu	Leu	Leu	Arg	Ile	Thr	His	Ser	Glu	Asp	Trp	Lys	Leu
	50					55					60				
Trp	Arg	Cys	Arg	Leu	Arg	Leu	Lys	Ser	Phe	Thr	Ser	Met	Asp	Ser	Arg
	65				70					75					80
Ser	Ala	Ser	His	Arg	Ser	Thr	Arg	Phe	Ala	Ala	Thr	Phe	Tyr	Asp	Ile
				85					90					95	
Glu	Thr	Leu	Lys	Val	Ile	Asp	Glu	Glu	Trp	Gln	Arg	Thr	Gln	Cys	Ser
			100					105					110		
Pro	Arg	Glu	Thr	Cys	Val	Glu	Val	Ala	Ser	Glu	Leu	Gly	Lys	Ser	Thr
		115					120					125			
sn	Thr	Phe	Phe	Lys	Pro	Pro	Cys	Val	Asn	Val	Phe	Arg	Cys	Gly	Gly
	130					135					140				
Cys	Cys	Asn	Glu	Glu	Ser	Leu	Ile	Cys	Met	Asn	Thr	Ser	Thr	Ser	Tyr
145					150					155					160
Ile	Ser	Lys	Gln	Leu	Phe	Glu	Ile	Ser	Val	Pro	Leu	Thr	Ser	Val	Pro
				165					170					175	
Glu	Leu	Val	Pro	Val	Lys	Val	Ala	Asn	His	Thr	Gly	Cys	Lys	Cys	Leu
			180					185					190		
Pro	Thr	Ala	Pro	Arg	His	Pro	Tyr	Ser	Ile	Ile	Arg	Arg	Ser	Ile	Gln
		195					200					205			
Ile	Pro	Glu	Glu	Asp	Arg	Cys	Ser	His	Ser	Lys	Lys	Leu	Cys	Pro	Ile
	210					215					220				
Asp	Met	Leu	Trp	Asp	Ser	Asn	Lys	Cys	Lys	Cys	Val	Leu	Gln	Glu	Glu
225					230					235					240
Asn	Pro	Leu	Ala	Gly	Thr	Glu	Asp	His	Ser	His	Leu	Gln	Glu	Pro	Ala
				245					250					255	
Leu	Cys	Gly	Pro	His	Met	Met	Phe	Asp	Glu	Asp	Arg	Cys	Glu	Cys	Val
			260					265					270		
Cys	Lys	Thr	Pro	Cys	Pro	Lys	Asp	Leu	Ile	Gln	His	Pro	Lys	Asn	Cys
		275					280					285			
Ser	Cys	Phe	Glu	Cys	Lys	Glu	Ser	Leu	Glu	Thr	Cys	Cys	Gln	Lys	His
	290					295					300				
Lys	Leu	Phe	His	Pro	Asp	Thr	Cys	Ser	Cys	Glu	Asp	Arg	Cys	Pro	Phe
305					310					315					320
His	Thr	Arg	Pro	Cys	Ala	Ser	Gly	Lys	Thr	Ala	Cys	Ala	Lys	His	Cys
				325					330					335	
Arg	Phe	Pro	Lys	Glu	Lys	Arg	Ala	Ala	Gln	Gly	Pro	His	Ser	Arg	Lys
			340					345					350		

Asn Pro

1. The first part of the document is a list of references. The references are listed in alphabetical order of the author's name. The references are as follows: